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GenCore version 5.1.6
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OM protein - protein search, using sw model

May 31, 2003, 17:57:32 ; Search time 15.6444 Seconds Run on:

(without alignments)
828.194 Million cell updates/sec

US-09-772-103-8

Perfect score:

1 MDFQVQIFSFLLISASVILS.........CQQWSSYPLTFGGGTKVBIK 128 Sequence:

Gapop 10.0 , Gapext 0.5 BLOSUM62 Scoring table:

383519 seqs, 101223694 residues cched:

'number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

| Cgn2_6/ptodata/1/pubpaa/USO8 NEW PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO8 NEW PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO6 NEW PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO6 PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO7_NEW PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO7_NEW PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO7_NEW PUB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
| Cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:* Published_Applications_AA:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Sequence 92, Appl Sequence 52, Appl Sequence 99, Appl Sequence 113, App Sequence 17, Appl Sequence 93, Appl Sequence 97, Appl Sequence 96, Appl Sequence 96, Appl Sequence 50, Appl Sequence 61, Appl Sequence 71, Appl Sequence 87, Appl Sequence 89, Appl Sequence 4, Appli Sequence 6, Appli Sequence 5, Appli Sequence 94, App. Sequence 111, Description 10 US-09-910-059-97 9 US-10-160-232-96 9 US-10-165-752-111 9 US-10-195-752-113 10 US-09-910-059-17 9 US-10-160-232-93 US-09-910-059-52 US-09-910-059-99 US-09-910-059-50 US-09-910-059-61 US-09-910-059-71 US-09-753-436-96 US-10-160-232-94 US-10-160-232-87 US-10-160-232-89 US-09-967-719C-6 US-09-795-515-5 Query Match Length DB Score Result 8 110 117 118 118 118

2333,	255 255 13,	Sequence 4, Appli Sequence 9, Appli Sequence 99, Appl Sequence 233, App Sequence 233, App Sequence 247, App Sequence 247, App Sequence 233, App Sequence 233, App Sequence 233, App Sequence 233, App
106 9 US-09-771-415-1 106 10 US-09-796-848A-3 213 9 US-09-996-288-231 213 9 US-09-996-265-231 106 9 US-09-996-288-54 1106 9 US-09-996-288-54 213 9 US-09-996-288-211	9 H H D D D D D D	128 9 US-09-905-928-4 128 9 US-10-096-964-4 129 10 US-10-965-099-99 213 9 US-09-996-288-233 213 9 US-09-996-288-247 213 9 US-09-996-288-247 213 9 US-09-996-288-247 213 9 US-09-996-288-247 213 9 US-09-996-288-247 213 9 US-09-996-265-239
20 487 74.4 22 487 74.4 23 487 74.4 24 487 74.4 25 488 73.9 26 484 73.9 27 484 73.9	4 682 73.7 4 882 73.7 4 882 73.6 6 881 73.6 6 881 73.4 7 881 73.4	, , , , , , , , , , , , , , , , , , ,
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ALIGNMENTS

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1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Synthetic protein
                                                                                                                                                                                                                 APPLICANT: KAWADA, YOKO
APPLICANT: NAKAMURA, KAZUYASU
APPLICANT: SIBUYA, MASABUMI
TITLE OF INVENTION: ANTI-HUMAN VEGF RECEPTOR FLT-1 MONOCLONAL ANTIBODY
FILE REFERENCE: 249-107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 91.4%; Pred. No. 2.7e-36;
Matches 117; Conservative 5; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                CURRENT FILING DATE: 2002-06-04
PRIOR APPLICATION UNDERS: US/09/453,718
PRIOR FILING DATE: 1999-12-03
PRIOR FILING DATE: 1999-12-03
PRIOR PAPLICATION NUMBER: 09/315,051
PRIOR APPLICATION NUMBER: 09/119,014
PRIOR PLILNG DATE: 1999-05-20
PRIOR PLILNG DATE: 1999-07-20
PRIOR PLILNG DATE: 1999-07-20
PRIOR PLILNG DATE: 1999-11-21
PRIOR FILING DATE: 1997-11-21
NUMBER OF SEQ ID NOS: 96
SOFTWARRE: PATCHILL VET: 2.1
                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/160,232 CURRENT FILING DATE: 2002-06-04
                                                      ; Sequence 92, Application US/10160232; Publication No. US20030088075A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                       APPLICANT: SHITARA, KENYA
                                                                                                                                                                 APPLICANT: ITO, MIKITO
APPLICANT: HANAI, NOBUO
APPLICANT: KAWADA, YOKO
RESULT 1
US-10-160-232-92
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61 PGKAPKLLIYDTSNLASGVPSRFSGSGGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120

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61 PGKAPKLLIYDTSNLASGVPSRFSGSGSGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
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APPLICANT: HANAL, NOBUO
APPLICANT: KAWADA, YOKU
APPLICANT: KAWADA, YOKU
APPLICANT: NAKAWURA, KAZUYASU
APPLICANT: NAKAWURA, MASABUMI
ITILE OF INVENTION: ANTI-HUMAN VEGF RECEPTOR FLT-1 MONOCLONAL ANTIBODY
FILE REFERENCE: 249-107
CURRENT APPLICATION NUMBER: 02/10/160,232
CURRENT FILING DATE: 2002-06-09
PRIOR PLING DATE: 1999-12-03
PRIOR PLING DATE: 1999-12-03
PRIOR PLING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: 09/119,014
PRIOR FILING DATE: 1999-07-20
PRIOR PLING DATE: 1999-11-21
NUMBER OF SEQ ID NOS: 96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 PGKAPKILLIYDTSNIASGVPSRFSGSGGTDYTLTISSLQPEDFATYXCQQWSSYPLTFG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 pakabkLwiystsniaksavbskresasasardvirtissiopediatyvoorstypiira 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: Synthetic protein
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                                                                                                                                                                                                                                                                            , OTHER INFORMATION: humanised light chain Fd sequence US-09-910-059-99
PRIOR APPLICATION NUMBER: GB 9703103.3 PRIOR FILING DATE: 1997-02-14 PRIOR APPLICATION NUMBER: GB9609405.7 PRIOR FILING DATE: 1996-05-04 NUMBER OF SEQ ID NOS: 131 SOFTWARE: PATENTIN VERSION 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 94, Application US/10160232; Publication No. US20030088075A1; GENERAL INFORMATION:
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Matches 116; Conservative
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Best Local Similarity
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SEQ ID NO 94
LENGTH: 126
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                                                                                                                                                               SEQ ID NO 99
LENGTH: 235
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APPLICANT: Copied, Michael Derek
APPLICANT: Copied, Michael Derek
APPLICANT: Edge, Michael Derek
APPLICANT: Edge, Michael Derek
APPLICANT: Edge, Michael Derek
APPLICANT: Emery. Stephen Charles
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody,
TITLE OF INVENTION: Their Therapeutic use in an Adept System
FILE REPRENCE: 1991-009
CURRENT APPLICATION NUMBER: US/09/910,059
CURRENT FILING DATE: 2001-07-23
PRIOR FILING DATE: 1998-10-29
PRIOR FILING DATE: PCT/GB97/01165
PRIOR FILING DATE: 1997-04-29
                                                                                                                                                                                                                                                              Sequence 52, Application US/09910059
Sequence 52, Application US/09910059
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Copley, Clive G
APPLICANT: Edge, Michael Derek
APPLICANT: Edge, Michael Derek
APPLICANT: Edge, Michael Derek
TITLE OF INVENTION: Their Therapeutic use in an Adept System
TITLE OF INVENTION: Their Therapeutic use in an Adept System
FILE REFERENCE: 1991-209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61 PGKAPKLLIYDISJULASGVPSRFSGSGSGTDYTLTISSLOPEDFATYYCQQWSSYPLTFG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
91.5%; Score 599; DB 10;
Best Local Similarity 91.4%; Pred. No. 5.5e-36;
Matches 117; Conservative 5; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TRIBENT APPLICATION NUMBER: US/09/910,059
CHRENT FILING DATE: 2001-07-23
CREAT FILING DATE: 2001-07-29
CREAT FILING DATE: 1998-10-29
PRIOR APPLICATION NUMBER: PCT/GB97/01165
PRIOR PILING DATE: 1997-04-29
PRIOR PLILING DATE: 1997-04-29
PRIOR PLILING DATE: 1997-02-14
PRIOR PILING DATE: 1997-02-14
PRIOR PILING DATE: 1996-05-04
NUMBER OF SEQ ID NOS: 131
SOFTWARE: PATENTIN VERSION 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 99, Application US/09910059
Patent No. US20020142359A1
GENERAL INFORMATION:
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                                                                                                      121 GGTKVEIK 128
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                                                                                                                                                                                                                                      RESULT 2
US-09-910-059-52
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LENGTH: 235
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61 PGKAPKLLIYDTSNLASGVPSRFSGSGSGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 PGKAPKLLIYRTSNLASGVPSRFSGSGGTFYTLTISSLQPEDFATYYCHQWSMY--TFG 118
                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic protein US-10-160-232-96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 87.5%; Score 573; DB 9; Length 126; Best Local Similarity 89.1%; Pred. No. 2.3e-34; Matches 114; Conservative 5; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: AKLINGIONA
COUNTRY: U.S.A.
ZIP: 22201-4714
COMPUTER READBELE FORM:
MEDIUM TYPE: Floppy disk
COMFUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
CLASSIFICATION: <Unknown>
PRIOR APPLICATION AUNGER: US/09/393,385B
FILING DATE: 27-JUN-96
TELECOMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KUWANA, YOSHIHISA
HASERAMA, MAMORU
TITLE OF INVENTION: HUMANIZED ANTIBODIES
NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
ADDRESSEE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD
              PRIOR APPLICATION NUMBER: 09/315,051
PRIOR FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: 09/119,014
PRIOR FILING DATE: 1998-07-20
PRIOR PRIOR DATE: 1998-07-20
PRIOR PRIING DATE: 1997-11-21
NUMBER OF SEQ ID NOS: 96
SEQ ID NO 96
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Publication No. US20030077276A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: NAKAMURA, KAZUYASU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (703)816-4000
TELEFAX: (703)816-4100
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SHITARA, KENYA
HANAI, NOBUO
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                  LENGTH:
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                                                                                                                                                                                                                                                                  ## Sequence 97, Application US/09910059
### APPLICANT: Copley, Clive G
### APPLICANT: Edge, Michael Derek
### APPLICANT: Sephen Charles
### TITLE OF INVENTION: Their Therapeutic use in an Adept System
### TITLE OF INVENTION: Their Therapeutic use in an Adept System
### APPLICATION NUMBER: US 09/171,945
### RENT FILING DATE: 1999-10-29
### PRIOR FILING DATE: 1999-10-29
### PRIOR PPLICATION NUMBER: GB 9703103.3
### PRIOR PLILING DATE: 1996-05-04
### PRIOR FILING DATE: DOWS: 131
### SOFTWARE: PATEUTING DATE: DOWS: 131
61 PGKAPKLLIYRTSNLASGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCHQWSMY--TFG 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 PGKAPKLLIYDTSNIASGVPSRFSGSGSGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61 PCKAPKLLIYSTSNLASGVPSRFSGSGSGTDYTFTISSLQPEDIATYYCQQRSTYPLTFG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-160-232-96

US-10-160-232-96

Sequence 96, Application US/10160232

Publication No. US20030088075A1

APPLICANT: SHITARA, KENYA

APPLICANT: TTO, MIKITO

APPLICANT: HANAI, NOBUO

APPLICANT: SHIBUTA, YOKO

APPLICANT: SHIBUTA, NASABUMI

TITLE OF INVENTION: ANTI-HUMAN VEGF RECEPTOR FLT-1 MONOCLONAL ANTIBODY

FILE REFERENCE: 249-107

CURRENT APPLICATION NUMBER: US/10/160,232

CURRENT FILING DATE: 2002-06-04

PRIOR FILING DATE: 1999-12-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK
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89.1%; Pred. No. 7.6e-35;
Live 6; Mismatches 8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: humanised light chain Fd sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 89.18
Matches 114; Conservative
                                                                                      121 GGTKVEIK 128
                                                                                                                                         119 QGTKVEIK 126
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121 QGTKLEIK 128
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SEQ ID NO 97
LENGTH: 235
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                                                                                                                                                                              1 MDFQVQIFSFLLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CUNTRY: U.S.A.

ZIP: 22201-4714

COUNTRY: U.S.A.

ZIP: 22201-4714

COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/195,752

FILING DATE: 16-Jul-2002

CLASSIFICATION: <UNKNOWN>
                                                                                                     Length 130;
                                                                                                                                             10; Indels
                                                                                                 Score 567; DB 9;
Pred. No. 6.2e-34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KUWANÀ, YOSHIHISA
HASEGANÀ, WANORU
TITLE OF INVENTION: HUMANIZED ANTIBODIES
NUMBER OF SEQUENCES: 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/393,385B
FILING DATE: 27-JUN-96
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)816-4000
TELEPRAX: (703)816-4100
                                                                                                                                             7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: NIXON & VANDERHYE P.C. STREET: 1100 NORTH GLEBE ROAD CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 113:
US-10-195-752-113
      MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 111:

US-10-195-752-111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 113, Application US/10195752
Publication No. US20030077276A1
GENERAL INFORMATION:
APPLICANT: NAKAMURA, KAZUYASU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 130 amino acids TYPE: amino acid
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SHITARA, KENYA
HANAI, NOBUO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 113:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
                                                                                                 Duery Match 86.6%; 8est Local Similarity 86.7%; 4atches 111; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 85.9
Matches 110; Conservative
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                           121 GGTKVEIK 128
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                                                                                                       Query Match
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1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQX 60

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APPLICANT: Copley, Clive G
APPLICANT: Edge, Michael Derek
APPLICANT: Edge, Michael Derek
APPLICANT: Emery, Stephen Charles
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody,
TITLE OF INVENTION: Their Therapeutic use in an Adept System
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 PGKAPKLLIYDTSNIASGVPSRFSGSGSGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 PGTSPKLWIYSTSNLASGVPARFSGSGGTSYSLTISRMEAEDAATYVCQQRSTYPLTFG 120
                                                       61 PGKAPKILLIYDTSNLASGVPSRFSGSGSGTDYTLTISSLOPEDFATYYCQQWSSYPLTFG 120
                                                                                    61 PGKAPKIMIYSTSNLASGVPARFSGSGSGTSYSLTISRLQPEDIATYYCQQRSSYPYTFG 120
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1 MHFQVQIFSFLLISASVIMSRGDIQLTQSPSSLSASPGDRVTITCSASSSVSYMHWFQQK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK
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APPLICANT: NAKAMURA, KAZUYASU
APPLICANT: SHIBUYA, MAZABUMI
ITILE OF INVENTION: ANTI-HUMAN VEGF RECEPTOR FLT-1 MONOCLONAL ANTIBODY
FILE REFERENCE: 249-107
CURRENT APPLICATION NUMBER: US/10/160,232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Pest Local Similarity 77.3%; Pred. No. 1.5e-30;
Matches 99; Conservative 17; Mismatches 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: chimaeric light chain sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 1991-209
CURRENT APPLICATION NUMBER: US/09/910,059
CURRENT FILING DATE: 2001-07-23
PRIOR APPLICATION NUMBER: US 09/171,945
PRIOR FILING DATE: 1998-10-29
PRIOR FILING DATE: 1998-10-29
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1996-05-04
NUMBER OF SEQ ID NOS: 131
SOFTWARE: Patentin version 3.1
LENGTH: 235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 93, Application US/10160232; Publication No. US20030088075A1; GENERAL INFORMATION: APPLICANT: SHITARA, KENYA
                                                                                                                                                                                                                                                                                                                                           ; Sequence 17, Application US/09910059; Patent No. US20020142359A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            121 GGTKVEIK 128
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61 PGKAPKLLIYDTSNLASGVPSRFSGSGGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: Synthetic protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ry Match 79.4%; Score 520; DB 9; Length 126; Local Similarity 79.7%; Pred. No. 1.4e-30; Ches 102; Conservative 9; Mismatches 15; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & Borun
STREET: 6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Gallarin, W. Michael
APPLICANT: Vazeux, Rosemay
TITLE OF INVENTION: ICAM-Related Materials and Methods
NUMBER OF SEQUENCES: 120
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 6300 Sears Tower, 233 South Wacker Drive CITY: Chicago Sears Tower, 233 South Wacker Drive CITY: Chicago Sears Tower, 233 South Wacker Drive CITY: Chicago States of America 21P: 60606-6402 COWPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk COMPUTER: 1BM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentl N Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/753,436
CURRENT FILING DATE: 2002-06-04

PRIOR APPLICATION NUMBER: US/09/453,718

PRIOR FILING DATE: 1999-12-03

PRIOR PELICATION NUMBER: 09/315,051

PRIOR PELICATION NUMBER: 09/115,051

PRIOR PELICATION NUMBER: 09/119,014

PRIOR PELING DATE: 1998-07-20

PRIOR PELING DATE: 1998-07-20

PRIOR PELING DATE: 1998-07-20

PRIOR PELING DATE: 1997-11-21

PRIOR FILING DATE: 1997-11-21

SEQ ID NO 93

LENGTHARE: PATCHITU Ver. 2.1

LENGTH: 126
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/286,754
FILING DATE: 05-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/102,852
FILING DATE: 05-AUG-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/487,113
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/382,289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 96, Application US/09753436; Patent No. US20010029293A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                121 GGTKVEIK 128
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                                                                                                                                                                                                                                                                                                                                                                                                                         ; US-10-160-232-93
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US-09-753-436-96
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Sequence 50, Application US/09910059
; Sequence 50, Application US/09910059
; Patent No. US20020142359al
; GENERAL INFORMATION:
    APPLICANT: Copley, Clive G
; APPLICANT: Edge, Michael Derek
    APPLICANT: Edge, Michael Derek
    APPLICANT: Edge, Michael Derek
    APPLICANT: Edge, Michael Derek
    APPLICANT: Edge, Michael Derek
    TITLE OF INVENTION: Their Therapeutic use in an Adept System
    TITLE OF INVENTION: Their Therapeutic use in an Adept System
    TILLE OF INVENTION: NUMBER: US/09/910,059
    CURRENT APPLICATION NUMBER: US 09/171,945
    PRIOR PILING DATE: 1998-10-29
    PRIOR FILING DATE: 1997-04-29
    PRIOR FILING DATE: 1997-04-29
    PRIOR PILING DATE: 1997-04-29
    PRIOR PILING DATE: 1997-02-14
    PRIOR PILING DATE: 1996-05-04
    PRIOR PILING DATE: 1996-05-04
    PRIOR PILING DATE: 1996-05-04
    PRIOR FILING DATE: 1996-05-04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 ARCDIQMTQSPSSLSASVGDRVTITCSASSSVSYIYWYQQKPGKAPKLLIYLTSNLASGV 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         62 PSRFSGSGSGTDFTLTISSLQPEDFATYYCQQWKSIPLTFGQGTKLEIK 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       76.3%; Score 500; DB 10; Length 110; 88.1%; Pred. No. 3.3e-29; tive 7; Mismatches 6; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 107;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      76.2%; Score 499; DB 10; 90.6%; Pred. No. 3.8e-29;
                  FILING DATE: 22-JAN-1993
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/894,061
FILING DATE: 05-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/827,689
FILING DATE: 26-MAY-1992
PRIOR APPLICATION NUMBER: US 07/827,689
FILING DATE: 27-JAN-1992
ATTORNEY/AGENT INFORMATION:
NAME: Williams, JOSEPh A., Jr.
REGISTRATION NUMBER: 33,282
TELECOMMUNICATION INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
   US 08/009,266
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TELEFAX: (312) 474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 96:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  110 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 88.1%
Matches 96; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear; MOLECULE TYPE: protein US-09-753-436-96
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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US-09-910-059-50
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61 PGKAPKILIYDTSNLASGVPSRFSGSGSGTDYTLTISSLQPEDFATYYCQQWSSYPLTFG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: Synthetic protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 MDFQVQIFSFLLISASVILSRGDIQMTQSPSSLSASVGDRVTITCSATSSITYMSWYQQK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: NAKAMURA, KAZUYASU
APPLICANT: SHIBUYA, MASABUMI
TITLE OF INVENTION: ANTI-HUMAN VEGF RECEPTOR FLT-1 MONOCLONAL ANTIBODY
FILE REFERENCE: 249-107
                                                                                                                                                                                                                                                 ; OTHER INFORMATION: humanised light chain variable region variant US-09-910-059-71
                                                                                                                                                                                                                                                                                                                                Score 493; DB 10; Length 107; Pred. No. 1e-28; 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 FSGSGSGTDYTFTISSLQPEDIATYXCQQRSTYPLTFGQGTKLEIK 106
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75.3%; Score 493; DB 9; Length 12
Best Local Similarity 72.7%; Pred. No. 1.2e-28;
Matches 93; Conservative 18; Mismatches 17; Indels
                                                                                                                                                                                                                                                                                                                                                                                    4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT PILING DATE: 249-110/
CURRENT PILING DATE: 2002-06-04
PRIOR APPLICATION NUMBER: US/10/160,232
CURRENT FILING DATE: 2002-06-04
PRIOR FILING DATE: 1999-12-03
PRIOR FILING DATE: 1999-05-20
PRIOR PILING DATE: 1999-06-20
PRIOR PILING DATE: 1999-06-20
PRIOR APPLICATION NUMBER: 09/119,014
PRIOR FILING DATE: 1999-07-20
PRIOR PILING DATE: 1999-11-21
NUMBER OF SEQ ID NOS: 96
SOFTWARE: Patentin Ver. 2.1
SOFTWARE: Patentin Ver. 2.1
PRIOR FILING DATE: 1997-02-14
PRIOR APPLICATION NUMBER: GB9609405.7
PRIOR FILING DATE: 1996-05-04
NUMBER OF SEQ ID NOS: 131
SOFTWARE: PatentIn version 3.1
SEQ ID NO 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 87, Application US/10160232;
Publication No. US2030088075A1
GENERAL INFORMATION:
APPLICANT: SHITARA, KENYA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
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89.6%;
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                            95; Conservative
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APPLICANT: HANAI, NOBUO
APPLICANT: KAWADA, YOKO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              121 GGTKVEIK 128
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                                                                                                                                                        LENGTH: 107
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Best Local S
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APPLICANT: Copiey, Clive G
APPLICANT: Copiey, Clive G
APPLICANT: Edge, Michael Derek
APPLICANT: Emery, Stephen Charles
APPLICANT: Emery, Stephen Charles
APPLICANT: Emery, Stephen Charles
TLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody,
TLE OF INVENTION: Their Therapeutic use in an Adept System
TLE OF INVENTION WUBBER: US/09/910,059
CURRENT APPLICATION NUMBER: US 09/171,945
PRIOR APPLICATION NUMBER: US 09/171,945
PRIOR FILING DATE: 1998-10-29
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1996-05-04
NUMBER OF SEQ ID NOS: 131
SOFTWARE: Patentin version 3.1
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; Patent No. US20020142359A1
; General INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TOTAL CAPIC COPICY, Clive G
APPLICANT: Edge, Michael Derek
APPLICANT: Emery, Stephen Charles
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody,
TITLE OF INVENTION: Their Therapeutic use in an Adept System
FILE REFERENCE: 1991-209
CURRENT APPLICATION NUMBER: US/09/910,059
CURRENT APPLICATION NUMBER: US/09/1171,945
PRIOR *TILING DATE: 1998-10-29
; PRIOR *TILING DATE: 1997-04-29
; PRIOR FILING DATE: 1997-04-29
; PRIOR FILING DATE: 1997-04-29
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                                                                                                         1 DIOMIOSPSSLSASVGDRVITICSASSSVTYMHWYQQKPGKAPKLLIYSTSNLASGVPSR 60
                                                         DIOMIQSPSSLSASVGDRVIITCSATSSITYMSWYQQKPGKAPKLLIYDTSNLASGVPSR 82
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          4; Mismatches
                                                                                                                                                                                                                                                                                                                                                Sequence 61, Application US/09910059 Patent No. US20020142359A1
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          96; Conservative
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LENGTH: 107
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Search completed: May 31, 2003, 18:05:42 Job time : 17.6444 secs

